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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2010; month=1; day=26; hr=13; min=32; sec=52; ms=661; ]

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\*\*\*\*\*

Reviewer Comments:

<210> 1  
<211> 405  
<212> DNA  
<213> Murine anti-B7-2 heavy chain

<220>

<221> CDS  
<222> (1)..(405)  
<223>

Numeric Identifier <213> can only be one of three choices, "Scientific name, i.e. Genus/species, Unknown or Artificial Sequence." For all sequences using "Unknown or Artificial sequence", for numeric identifier <213>, a mandatory feature is required to explain the source of the genetic material. The feature consists of <220>, which remains blank, and <223>, which states the source of the genetic material. Suggest using "Artificial sequence" for numeric identifier <213> and "The above given response" for numeric identifier <223> in the mandatory feature. Please check for similar errors and make all necessary changes.

\*\*\*\*\*

Application No: 09501102 Version No: 2.0

**Input Set:**

**Output Set:**

**Started:** 2010-01-25 14:23:49.649  
**Finished:** 2010-01-25 14:23:56.105  
**Elapsed:** 0 hr(s) 0 min(s) 6 sec(s) 456 ms  
**Total Warnings:** 44  
**Total Errors:** 31  
**No. of SeqIDs Defined:** 52  
**Actual SeqID Count:** 52

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
E 201	Mandatory field data missing in <223> in SEQ ID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
E 201	Mandatory field data missing in <223> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (5)
E 201	Mandatory field data missing in <223> in SEQ ID (5)
W 402	Undefined organism found in <213> in SEQ ID (6)
W 402	Undefined organism found in <213> in SEQ ID (7)
E 201	Mandatory field data missing in <223> in SEQ ID (7)
W 402	Undefined organism found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)

### Input Set:

### **Output Set:**

**Started:** 2010-01-25 14:23:49.649  
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**Elapsed:** 0 hr(s) 0 min(s) 6 sec(s) 456 ms  
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**Input Set:**

**Output Set:**

**Started:** 2010-01-25 14:23:49.649  
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**Elapsed:** 0 hr(s) 0 min(s) 6 sec(s) 456 ms  
**Total Warnings:** 44  
**Total Errors:** 31  
**No. of SeqIDs Defined:** 52  
**Actual SeqID Count:** 52

Error code	Error Description
E 336	Empty lines found between the proteins and the dna
E 336	Empty lines found between the proteins and the dna
E 336	Empty lines found between the proteins and the dna
W 402	Undefined organism found in <213> in SEQ ID (24)
W 213	Artificial or Unknown found in <213> in SEQ ID (25)
W 213	Artificial or Unknown found in <213> in SEQ ID (26)
W 213	Artificial or Unknown found in <213> in SEQ ID (27)
W 213	Artificial or Unknown found in <213> in SEQ ID (28)
W 213	Artificial or Unknown found in <213> in SEQ ID (29)
W 213	Artificial or Unknown found in <213> in SEQ ID (30)
W 213	Artificial or Unknown found in <213> in SEQ ID (31)
W 213	Artificial or Unknown found in <213> in SEQ ID (32) This error has occurred more than 20 times, will not be displayed
W 402	Undefined organism found in <213> in SEQ ID (41)
E 322	CDS location out of range SEQID (41) At Protein count (133)
E 320	Wrong Nucleic Acid Designator, ga in SEQID (41)
W 402	Undefined organism found in <213> in SEQ ID (42)
W 402	Undefined organism found in <213> in SEQ ID (43)
E 322	CDS location out of range SEQID (43) At Protein count (136)
E 320	Wrong Nucleic Acid Designator, cc in SEQID (43)
E 322	CDS location out of range SEQID (43) At Protein count (234)
E 320	Wrong Nucleic Acid Designator, ag in SEQID (43)
E 322	CDS location out of range SEQID (43) At Protein count (246)

**Input Set:**

**Output Set:**

**Started:** 2010-01-25 14:23:49.649  
**Finished:** 2010-01-25 14:23:56.105  
**Elapsed:** 0 hr(s) 0 min(s) 6 sec(s) 456 ms  
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**Total Errors:** 31  
**No. of SeqIDs Defined:** 52  
**Actual SeqID Count:** 52

Error code	Error Description
E 320	Wrong Nucleic Acid Designator, ca in SEQID (43)
E 322	CDS location out of range SEQID (43) At Protien count (355)
E 320	Wrong Nucleic Acid Designator, gg in SEQID (43)
W 402	Undefined organism found in <213> in SEQ ID (44)

SEQUENCE LISTING

<110> Co, Man Sung  
Vasquez, Maximiliano  
Carreno, Beatriz  
Celniker, Abbie Cheryl  
Collins, Mary  
Goldman, Samuel  
Gray, Gary S.  
Knight, Andrea  
O'Hara, Denise  
Rup, Bonita  
Veldman, Geertruida M.

<120> HUMANIZED IMMUNOGLOBULIN REACTIVE WITH B7-2 MOLECULES AND METHODS OF TREATMENT THEREWITH

<130> 08702.0081-00000

<140> 09501102  
<141> 2000-02-09

<150> 09/249,011  
<151> 1999-02-12

<160> 52

<170> PatentIn version 3.1

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<212> DNA  
<213> Murine anti-B7-2 heavy chain

<220>

<221> CDS  
<222> (1)..(405)  
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1 5 10 15

gtg cac tcc cag gtc cag ctg cag tct ggg cct gag ctg gtg agg 96  
Val His Ser Gln Val Gln Leu Gln Ser Gly Pro Glu Leu Val Arg  
20 25 30

cct ggg gaa tca gtg aag att tcc tgc aag ggt tcc ggc tac aca ttc 144  
Pro Gly Glu Ser Val Lys Ile Ser Cys Lys Gly Ser Gly Tyr Thr Phe  
35 40 45

act gat tat gct ata cag tgg gtg aag cag agt cat gca aag agt cta 192  
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50 55 60

gag tgg att gga gtt att aat att tac tat gat aat aca aac tac aac 240  
Glu Trp Ile Gly Val Ile Asn Ile Tyr Tyr Asp Asn Thr Asn Tyr Asn

65 70 75 80

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Gln Lys Phe Lys Gly Lys Ala Thr Met Thr Val Asp Lys Ser Ser Ser  
85 90 95

aca gcc tat atg gaa ctt gcc aga ttg aca tct gag gat tct gcc atc 336  
Thr Ala Tyr Met Glu Leu Ala Arg Leu Thr Ser Glu Asp Ser Ala Ile  
100 105 110

tat tac tgt gca aga gcg gcc tgg tat atg gac tac tgg ggt caa gga 384  
Tyr Tyr Cys Ala Arg Ala Ala Trp Tyr Met Asp Tyr Trp Gly Gln Gly  
115 120 125

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Thr Ser Val Thr Val Ser Ser  
130 135

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<212> PRT  
<213> Murine anti-B7-2 heavy chain

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1 5 10 15

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20 25 30

Pro Gly Glu Ser Val Lys Ile Ser Cys Lys Gly Ser Gly Tyr Thr Phe  
35 40 45

Thr Asp Tyr Ala Ile Gln Trp Val Lys Gln Ser His Ala Lys Ser Leu  
50 55 60

Glu Trp Ile Gly Val Ile Asn Ile Tyr Tyr Asp Asn Thr Asn Tyr Asn  
65 70 75 80

Gln Lys Phe Lys Gly Lys Ala Thr Met Thr Val Asp Lys Ser Ser Ser  
85 90 95

Thr Ala Tyr Met Glu Leu Ala Arg Leu Thr Ser Glu Asp Ser Ala Ile  
100 105 110

Tyr Tyr Cys Ala Arg Ala Ala Trp Tyr Met Asp Tyr Trp Gly Gln Gly  
115 120 125

Thr Ser Val Thr Val Ser Ser  
 130 135

<210> 3  
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 <213> Murine anti-B7-2 light chain

<220>  
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 <222> (1)..(396)  
 <223>

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Met	Asp	Ser	Gln	Ala	Gln	Val	Leu	Ile	Leu	Leu	Leu	Leu	Trp	Val	Ser		
1		5						10						15			

ggt	acc	tgt	ggg	gac	att	gtg	ctg	tca	cag	tct	cca	tcc	tcc	ctg	gct		96
Gly	Thr	Cys	Gly	Asp	Ile	Val	Leu	Ser	Gln	Ser	Pro	Ser	Ser	Leu	Ala		
20		25												30			

gtg	tca	gca	gga	gag	aag	gtc	act	atg	agc	tgc	aaa	tcc	agt	cag	agt		144
Val	Ser	Ala	Gly	Glu	Lys	Val	Thr	Met	Ser	Cys	Lys	Ser	Ser	Gln	Ser		
35		40												45			

ctg	ctc	aac	agt	aga	acc	cga	gag	aac	tac	ttg	gct	tgg	tac	cag	cag		192
Leu	Leu	Asn	Ser	Arg	Thr	Arg	Glu	Asn	Tyr	Leu	Ala	Trp	Tyr	Gln	Gln		
50		55												60			

aaa	cca	ggg	cag	tct	cct	aaa	ctg	ctg	atc	tac	tgg	gca	tcc	act	agg		240
Lys	Pro	Gly	Gln	Ser	Pro	Lys	Leu	Leu	Ile	Tyr	Trp	Ala	Ser	Thr	Arg		
65		70												80			

gaa	tct	ggg	gtc	cct	gat	cgc	ttc	aca	ggc	agt	gga	tct	ggg	aca	gat		288
Glu	Ser	Gly	Val	Pro	Asp	Arg	Phe	Thr	Gly	Ser	Gly	Ser	Gly	Thr	Asp		
85		90												95			

ttc	act	ctc	acc	atc	agc	agt	gtg	cag	gct	gaa	gac	ctg	gca	gtt	tat		336
Phe	Thr	Leu	Thr	Ile	Ser	Ser	Val	Gln	Ala	Glu	Asp	Leu	Ala	Val	Tyr		
100		105												110			

tac	tgc	acg	caa	tct	tat	aat	ctt	tac	acg	ttc	gga	ggg	ggg	acc	aag		384
Tyr	Cys	Thr	Gln	Ser	Tyr	Asn	Leu	Tyr	Thr	Phe	Gly	Gly	Gly	Thr	Lys		
115		120												125			

ctg	gaa	ata	aaa													396
Leu	Glu	Ile	Lys													
130																

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 <213> Murine anti-B7-2 light chain

<400> 4

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Val Ser Ala Gly Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser  
35 40 45

Leu Leu Asn Ser Arg Thr Arg Glu Asn Tyr Leu Ala Trp Tyr Gln Gln  
50 55 60

Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg  
65 70 75 80

Glu Ser Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp  
85 90 95

Phe Thr Leu Thr Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Val Tyr  
100 105 110

Tyr Cys Thr Gln Ser Tyr Asn Leu Tyr Thr Phe Gly Gly Thr Lys  
115 120 125

Leu Glu Ile Lys  
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<210> 5  
<211> 405  
<212> DNA  
<213> Humanized murine anti-human B7-2 heavy chain

<220>  
<221> CDS  
<222> (1)...(405)  
<223>

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1 5 10 15

gtg cac tcc cag gtc cag ctg gtg cag tct ggg gct gag gtg aag aag 96  
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys  
20 25 30

cct	ggg	agc	tca	gtg	aag	gtg	tcc	tgc	aaa	gct	tcc	ggc	tac	aca	ttc		144	
Pro	Gly	Ser	Ser	Val	Lys	Val	Ser	Cys	Lys	Ala	Ser	Gly	Tyr	Thr	Phe			
35																45		
act	gat	tat	gct	ata	cag	tgg	gtg	aga	cag	gct	cct	gga	cag	ggc	ctc		192	
Thr	Asp	Tyr	Ala	Ile	Gln	Trp	Val	Arg	Gln	Ala	Pro	Gly	Gln	Gly	Leu			
50																60		
gag	tgg	att	gga	gtt	att	aat	att	taa	tat	gat	aat	aca	aac	taa			240	
Glu	Trp	Ile	Gly	Val	Ile	Asn	Ile	Tyr	Tyr	Asp	Asn	Thr	Asn	Tyr	Asn			
65																80		
cag	aag	ttt	aag	ggc	aag	gcc	aca	atg	act	gta	gac	aag	tcg	acg	acg		288	
Gln	Lys	Phe	Lys	Gly	Lys	Ala	Thr	Met	Thr	Val	Asp	Lys	Ser	Thr	Ser			
85																95		
aca	gcc	tat	atg	gaa	ctt	agt	tct	ttg	aga	tct	gag	gat	acg	gcc	gtt		336	
Thr	Ala	Tyr	Met	Glu	Leu	Ser	Ser	Leu	Arg	Ser	Glu	Asp	Thr	Ala	Val			
100																110		
tat	tac	tgt	gca	aga	gca	gag	gcc	tgg	tat	atg	gac	tac	tgg	ggt	caa	ggt		384
Tyr	Tyr	Cys	Ala	Arg	Ala	Ala	Trp	Tyr	Met	Asp	Tyr	Tyr	Trp	Gly	Gln	Gly		
115																125		
acc	ctt	gtc	acc	gtc	tcc	tca											405	
Thr	Leu	Val	Thr	Val	Ser	Ser												
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<213>	Humanized murine anti-human B7-2 heavy chain																	
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1																15		
Val	His	Ser	Gln	Val	Gln	Leu	Val	Gln	Ser	Gly	Ala	Glu	Val	Lys	Lys			
																20	25	30
Pro	Gly	Ser	Ser	Val	Lys	Val	Ser	Cys	Lys	Ala	Ser	Gly	Tyr	Thr	Phe			
35																45		
Thr	Asp	Tyr	Ala	Ile	Gln	Trp	Val	Arg	Gln	Ala	Pro	Gly	Gln	Gly	Leu			
50																60		
Glu	Trp	Ile	Gly	Val	Ile	Asn	Ile	Tyr	Tyr	Asp	Asn	Thr	Asn	Tyr	Asn			
65																75	80	

Gln Lys Phe Lys Gly Lys Ala Thr Met Thr Val Asp Lys Ser Thr Ser  
85 90 95

Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val  
100 105 110

Tyr Tyr Cys Ala Arg Ala Ala Trp Tyr Met Asp Tyr Trp Gly Gln Gly  
115 120 125

Thr Leu Val Thr Val Ser Ser  
130 135

<210> 7  
<211> 396  
<212> DNA  
<213> Humanized murine anti-human B7-2 light chain

<220>  
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<222> (1)...(396)  
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1 5 10 15

ggc acc tgt ggg gac att gtg ctg aca cag tct cca gat tcc ctg gct 96  
Gly Thr Cys Gly Asp Ile Val Leu Thr Gln Ser Pro Asp Ser Leu Ala  
20 25 30

gta agc tta gga gag agg gcc act att agc tgc aaa tcc agt cag agt 144  
Val Ser Leu Gly Glu Arg Ala Thr Ile Ser Cys Lys Ser Ser Gln Ser  
35 40 45

ctg ctc aac agt aga acc cga gag aac tac ttg gct tgg tac cag cag 192  
Leu Leu Asn Ser Arg Thr Arg Glu Asn Tyr Leu Ala Trp Tyr Gln Gln  
50 55 60

aaa cca ggg cag cct cct aaa ctg ctg atc tac tgg gca tcc act agg 240  
Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg  
65 70 75 80

gaa tct ggg gtc cct gat cgc ttc agt ggc agt gga tct ggg aca gat 288  
Glu Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp  
85 90 95

ttc act ctc acc atc agc agt ctg cag gct gaa gac gtg gca gtt tat 336  
Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr  
100 105 110

tac tgc acg caa tct tat aat ctt tac acg ttc gga cag ggg acc aag 384  
Tyr Cys Thr Gln Ser Tyr Asn Leu Tyr Thr Phe Gly Gln Gly Thr Lys

115

120

125

396

gtg gaa ata aaa  
Val Glu Ile Lys  
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<212> PRT  
<213> Humanized murine anti-human B7-2 light chain

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Gly Thr Cys Gly Asp Ile Val Leu Thr Gln Ser Pro Asp Ser Leu Ala  
20 25 30

Val Ser Leu Gly Glu Arg Ala Thr Ile Ser Cys Lys Ser Ser Gln Ser  
35 40 45

Leu Leu Asn Ser Arg Thr Arg Glu Asn Tyr Leu Ala Trp Tyr Gln Gln  
50 55 60

Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg  
65 70 75 80

Glu Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Thr Asp  
85 90 95

Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr  
100 105 110

Tyr Cys Thr Gln Ser Tyr Asn Leu Tyr Thr Phe Gly Gln Gly Thr Lys  
115 120 125

Val Glu Ile Lys  
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1 5 10 15

Gly

<210> 13  
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<223> CDR3 of humanized murine anti-human B7-2 heavy chain

<221> CDS  
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1 5 10 15

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1 5 10 15

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<400> 17  
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<400> 18

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<221> CDS

<222> (1) .. (24)

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